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## SEQUENCE LISTING

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<110> Myers, Lisa E.
     Schryvres, Anthony B.
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     Loosmore, Sheena M.
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     Yang, Yan-Ping
     Klein, Michel H.
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Asn Leu Val Val Val Leu Asp Glu Thr Val Val Thr Ala Lys Lys Asn 50 55 60

Ala Arg Lys Ala Asn Glu Val Thr Gly Leu Gly Lys Val Val Lys Thr 65 70 75 80

Ala Glu Thr Ile Asn Lys Glu Gln Val Leu Asn Ile Arg Asp Leu Thr 85 90 95

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Val Asp Gly Ile Asn Gln Ala Gln His Tyr Ala Leu Gln Gly Pro Val 130 135 140

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Lys Gly Thr Asp Asn Ile Tyr Gln Pro Asn Gln Ala Thr Val Val Lys 675 . 680 685

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Phe Ser Glu Met Tyr Gly Glu Arg Phe Gly Val Thr Ile Gly Lys Gly 785 790 795 800

Thr Gln His Gly Cys Lys Gly Leu Tyr Tyr Ile Cys Gln Gln Thr Val 805 810 815

His Gln Thr Lys Leu Lys Pro Glu Lys Ser Phe Asn Gln Glu Ile Gly 820 825 830

Ala Thr Leu His Asn His Leu Gly Ser Leu Glu Val Ser Tyr Phe Lys 835 840 845

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Gly Phe His Asn Gly Gln Asp Ala Asp Leu Thr Gly Ile Asn Ile Leu 885 890 895

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<212> PRT

<213> Moraxella catarrhalis

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Glu Val Thr Gly Leu Gly Lys Val Val Lys Thr Ala Glu Thr Ile Asn 50 55 60

Lys Glu Gln Val Leu Asn Ile Arg Asp Leu Thr Arg Tyr Asp Pro Gly 65 70 75 80

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Arg Gly Met Asp Lys Asn Arg Val Ala Val Leu Val Asp Gly Ile Asn 100 105 110

Gln Ala Gln His Tyr Ala Leu Gln Gly Pro Val Ala Gly Lys Asn Tyr 115 120 125

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Val Glu Ile Ser Lys Gly Ala Asn Ser Ser Glu Tyr Gly Ser Gly Ala 145 150 155 160

Leu Ser Gly Ser Val Ala Phe Val Thr Lys Thr Ala Asp Asp Ile Ile 165 170 175

Lys Asp Gly Lys Asp Trp Gly Val Gln Thr Lys Thr Ala Tyr Ala Ser 180 185 190

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Tyr Thr Gly Pro Asn Arg Leu Ile Pro Asn Pro Leu Thr Gln Asp Ser 290 295 300

Lys Ser Leu Leu Arg Pro Gly Tyr Gln Leu Asn Asp Lys His Tyr 315 Val Gly Gly Val Tyr Glu Ile Thr Lys Gln Asn Tyr Ala Met Gln Asp Lys Thr Val Pro Ala Tyr Leu Thr Val His Asp Ile Glu Lys Ser Arg Leu Ser Asn His Ala Gln Ala Asn Gly Tyr Tyr Gln Gly Asn Asn Leu 360 Gly Glu Arg Ile Arg Asp Thr Ile Gly Pro Asp Ser Gly Tyr Gly Ile Asn Tyr Ala His Gly Val Phe Tyr Asp Glu Lys His Gln Lys Asp Arg Leu Gly Leu Glu Tyr Val Tyr Asp Ser Lys Gly Glu Asn Lys Trp Phe Asp Asp Val Arg Val Ser Tyr Asp Lys Gln Asp Ile Thr Leu Arg Ser 425 Gln Leu Thr Asn Thr His Cys Ser Thr Tyr Pro His Ile Asp Lys Asn 435 440 Cys Thr Pro Asp Val Asn Lys Pro Phe Ser Val Lys Glu Val Asp Asn Asn Ala Tyr Lys Glu Gln His Asn Leu Ile Lys Ala Val Phe Asn Lys 475 Lys Met Ala Leu Gly Ser Thr His His His Ile Asn Leu Gln Val Gly Tyr Asp Lys Phe Asn Ser Ser Leu Ser Arg Glu Asp Tyr Arg Leu Ala Thr His Gln Ser Tyr Glu Lys Leu Asp Tyr Thr Pro Pro Ser Asn Pro Leu Pro Asp Lys Phe Lys Pro Ile Leu Gly Ser Asn Asn Lys Pro Ile Cys Leu Asp Ala Tyr Gly Tyr Gly His Asp His Pro Gln Ala Cys Asn Ala Lys Asn Ser Thr Tyr Gln Asn Phe Ala Ile Lys Lys Gly Ile Glu Gln Tyr Asn Gln Lys Thr Asn Thr Asp Lys Ile Asp Tyr Gln Ala Ile 580 Ile Asp Gln Tyr Asp Lys Gln Asn Pro Asn Ser Thr Leu Lys Pro Phe 600 Glu Lys Ile Lys Gln Ser Leu Gly Gln Glu Lys Tyr Asn Lys Ile Asp 610 615 620

Glu Leu Gly Phe Lys Ala Tyr Lys Asp Leu Arg Asn Glu Trp Ala Gly Trp Thr Asn Asp Asn Ser Gln Gln Asn Ala Asn Lys Gly Thr Asp Asn 650 Ile Tyr Gln Pro Asn Gln Ala Thr Val Val Lys Asp Asp Lys Cys Lys 665 Tyr Ser Glu Thr Asn Ser Tyr Ala Asp Cys Ser Thr Thr Ala His Ile 680 Ser Gly Asp Asn Tyr Phe Ile Ala Leu Lys Asp Asn Met Thr Ile Asn 695 Lys Tyr Val Asp Leu Gly Leu Gly Ala Arg Tyr Asp Arg Ile Lys His Lys Ser Asp Val Pro Leu Val Asp Asn Ser Ala Ser Asn Gln Leu Ser Trp Asn Phe Gly Val Val Lys Pro Thr Asn Trp Leu Asp Ile Ala 745 Tyr Arg Ser Ser Gln Gly Phe Arg Met Pro Ser Phe Ser Glu Met Tyr 760 Gly Glu Arg Phe Gly Val Thr Ile Gly Lys Gly Thr Gln His Gly Cys 775 Lys Gly Leu Tyr Tyr Ile Cys Gln Gln Thr Val His Gln Thr Lys Leu Lys Pro Glu Lys Ser Phe Asn Gln Glu Ile Gly Ala Thr Leu His Asn His Leu Gly Ser Leu Glu Val Ser Tyr Phe Lys Asn Arg Tyr Thr Asp Leu Ile Val Gly Lys Ser Glu Glu Ile Arg Thr Leu Thr Gln Gly Asp Asn Ala Gly Lys Gln Arg Gly Lys Gly Asp Leu Gly Phe His Asn Gly 855 Gln Asp Ala Asp Leu Thr Gly Ile Asn Ile Leu Gly Arg Leu Asp Leu Asn Ala Val Asn Ser Arg Leu Pro Tyr Gly Leu Tyr Ser Thr Leu Ala 885 Tyr Asn Lys Val Asp Val Lys Gly Lys Thr Leu Asn Pro Thr Leu Ala Gly Thr Asn Ile Leu Phe Asp Ala Ile Gln Pro Ser Arg Tyr Val Val 920 925

Gly Leu Gly Tyr Asp Ala Pro Ser Gln Lys Trp Gly Ala Asn Ala Ile 930 . 935 940

Phe Thr His Ser Asp Ala Lys Asn Pro Ser Glu Leu Leu Ala Asp Lys 945 950 955 960

Asn Leu Gly Asn Gly Asn Ile Gln Thr Lys Gln Ala Thr Lys Ala Lys 965 970 975

Ser Thr Pro Trp Gln Thr Leu Asp Leu Ser Gly Tyr Val Asn Ile Lys 980 985 990

Asp Asn Phe Thr Leu Arg Ala Gly Val Tyr Asn Val Phe Asn Thr Tyr 995 1000 1005

Tyr Thr Thr Trp Glu Ala Leu Arg Gln Thr Ala Glu Gly Ala Val Asn 1010 1015 1020

Gln His Thr Gly Leu Ser Gln Asp Lys His Tyr Gly Arg Tyr Ala Ala 1025 1030 1035 1040

Pro Gly Arg Asn Tyr Gln Leu Ala Leu Glu Met Lys Phe 1045

<210> 11

<211> 702

<212> PRT

<213> Moraxella catarrhalis

<400> 11

Met Lys His Ile Pro Leu Thr Thr Leu Cys Val Ala Ile Ser Ala Val 1 5 10 15

Leu Leu Thr Ala Cys Gly Gly Ser Gly Gly Ser Asn Pro Pro Ala Pro 20 25 30

Thr Pro Ile Pro Asn Ala Ser Gly Ser Gly Asn Thr Gly Asn Thr Gly 35 40 45

Asn Ala Gly Gly Thr Asp Asn Thr Ala Asn Ala Gly Asn Thr Gly Gly 50 55 60

Thr Asn Ser Gly Thr Gly Ser Ala Asn Thr Pro Glu Pro Lys Tyr Gln 65 70 75 80

Asp Val Pro Thr Glu Lys Asn Glu Lys Asp Lys Val Ser Ser Ile Gln 85 90 95

Glu Pro Ala Met Gly Tyr Gly Met Ala Leu Ser Lys Ile Asn Leu His

Asn Arg Gln Asp Thr Pro Leu Asp Glu Lys Asn Ile Ile Thr Leu Asp

Gly Lys Lys Gln Val Ala Glu Gly Lys Lys Ser Pro Leu Pro Phe Ser 130 135 140

Leu Asp Val Glu Asn Lys Leu Leu Asp Gly Tyr Ile Ala Lys Met Asn 150 155 Val Ala Asp Lys Asn Ala Ile Gly Asp Arg Ile Lys Lys Gly Asn Lys Glu Ile Ser Asp Glu Glu Leu Ala Lys Gln Ile Lys Glu Ala Val Arg 185 Lys Ser His Glu Phe Gln Gln Val Leu Ser Ser Leu Glu Asn Lys Ile 200 Phe His Ser Asn Asp Gly Thr Thr Lys Ala Thr Thr Arg Asp Leu Lys 215 Tyr Val Asp Tyr Gly Tyr Tyr Leu Ala Asn Asp Gly Asn Tyr Leu Thr Val Lys Thr Asp Lys Leu Trp Asn Leu Gly Pro Val Gly Gly Val Phe Tyr Asn Gly Thr Thr Ala Lys Glu Leu Pro Thr Gln Asp Ala Val 265 Lys Tyr Lys Gly His Trp Asp Phe Met Thr Asp Val Ala Asn Arg Arg Asn Arg Phe Ser Glu Val Lys Glu Asn Ser Gln Ala Gly Trp Tyr Tyr 295 Gly Ala Ser Ser Lys Asp Glu Tyr Asn Arg Leu Leu Thr Lys Glu Asp Ser Ala Pro Asp Gly His Ser Gly Glu Tyr Gly His Ser Ser Glu Phe Thr Val Asn Phe Lys Glu Lys Lys Leu Thr Gly Lys Leu Phe Ser Asn Leu Gln Asp Arg His Lys Gly Asn Val Thr Lys Thr Glu Arg Tyr Asp Ile Asp Ala Asn Ile His Gly Asn Arg Phe Arg Gly Ser Ala Thr Ala 375 Ser Asn Lys Asn Asp Thr Ser Lys His Pro Phe Thr Ser Asp Ala Asn Asn Arg Leu Glu Gly Gly Phe Tyr Gly Pro Lys Gly Glu Glu Leu Ala 405 410 Gly Lys Phe Leu Thr Asn Asp Asn Lys Leu Phe Gly Val Phe Gly Ala Lys Arg Glu Ser Lys Ala Glu Glu Lys Thr Glu Ala Ile Leu Asp Ala 440

Tyr Ala Leu Gly Thr Phe Asn Thr Ser Asn Ala Thr Thr Phe Thr Pro 450 455 460

Phe Thr Glu Lys Gln Leu Asp Asn Phe Gly Asn Ala Lys Lys Leu Val 465 470 475 480

Leu Gly Ser Thr Val Ile Asp Leu Val Pro Thr Asp Ala Thr Lys Asn 485 490 495

Glu Phe Thr Lys Asp Lys Pro Glu Ser Ala Thr Asn Glu Ala Gly Glu 500 505 510

Thr Leu Met Val Asn Asp Glu Val Ser Val Lys Thr Tyr Gly Lys Asn 515 520 525

Phe Glu Tyr Leu Lys Phe Gly Glu Leu Ser Ile Gly Gly Ser His Ser 530 540

Val Phe Leu Gln Gly Glu Arg Thr Ala Thr Thr Gly Glu Lys Ala Val 545 550 555 560

Pro Thr Thr Gly Thr Ala Lys Tyr Leu Gly Asn Trp Val Gly Tyr Ile 565 570 575

Thr Gly Lys Asp Thr Gly Thr Gly Thr Gly Lys Ser Phe Thr Asp Ala 580 585 590

Gln Asp Val Ala Asp Phe Asp Ile Asp Phe Gly Asn Lys Ser Val Ser 595 600 605

Gly Lys Leu Ile Thr Lys Gly Arg Gln Asp Pro Val Phe Ser Ile Thr 610 620

Gly Gln Ile Ala Gly Asn Gly Trp Thr Gly Thr Ala Ser Thr Thr Lys 625 630 635 640

Ala Asp Ala Gly Gly Tyr Lys Ile Asp Ser Ser Ser Thr Gly Lys Ser 645 650 655

Ile Ala Ile Lys Asp Ala Asn Val Thr Gly Gly Phe Tyr Gly Pro Asn 660 . 670

Ala Asn Glu Met Gly Gly Ser Phe Thr His Asn Ala Asp Asp Ser Lys 675 680 685

Ala Ser Val Val Phe Gly Thr Lys Arg Gln Gln Glu Val Lys 690 695 700

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<213> Moraxella catarrhalis

<400> 12

Cys Gly Gly Ser Gly Gly Ser Asn Pro Pro Ala Pro Thr Pro Ile Pro 1 5 10 15

Asn Ala Ser Gly Ser Gly Asn Thr Gly Asn Thr Gly Asn Ala Gly Gly 20 25 30

Thr Asp Asn Thr Ala Asn Ala Gly Asn Thr Gly Gly Thr Asn Ser Gly 35 40 45

Thr Gly Ser Ala Asn Thr Pro Glu Pro Lys Tyr Gln Asp Val Pro Thr 50 55 60

Glu Lys Asn Glu Lys Asp Lys Val Ser Ser Ile Gln Glu Pro Ala Met 65 70 75 80

Gly Tyr Gly Met Ala Leu Ser Lys Ile Asn Leu His Asn Arg Gln Asp 85 90 95

Thr Pro Leu Asp Glu Lys Asn Ile Ile Thr Leu Asp Gly Lys Lys Gln
100 105 110

Val Ala Glu Gly Lys Lys Ser Pro Leu Pro Phe Ser Leu Asp Val Glu 115 120 125

Asn Lys Leu Leu Asp Gly Tyr Ile Ala Lys Met Asn Val Ala Asp Lys 130 135 140

Asn Ala Ile Gly Asp Arg Ile Lys Lys Gly Asn Lys Glu Ile Ser Asp 145 150 155 160

Glu Glu Leu Ala Lys Gln Ile Lys Glu Ala Val Arg Lys Ser His Glu 165 170 175

Phe Gln Gln Val Leu Ser Ser Leu Glu Asn Lys Ile Phe His Ser Asn 180 185 190

Asp Gly Thr Thr Lys Ala Thr Thr Arg Asp Leu Lys Tyr Val Asp Tyr 195 200 205

Gly Tyr Tyr Leu Ala Asn Asp Gly Asn Tyr Leu Thr Val Lys Thr Asp 210 220

Lys Leu Trp Asn Leu Gly Pro Val Gly Gly Val Phe Tyr Asn Gly Thr 225 230 235 240

Thr Thr Ala Lys Glu Leu Pro Thr Gln Asp Ala Val Lys Tyr Lys Gly
245 250 255

His Trp Asp Phe Met Thr Asp Val Ala Asn Arg Arg Asn Arg Phe Ser 260 265 270

Glu Val Lys Glu Asn Ser Gln Ala Gly Trp Tyr Tyr Gly Ala Ser Ser 275 280 285

Lys Asp Glu Tyr Asn Arg Leu Leu Thr Lys Glu Asp Ser Ala Pro Asp 290 295 300

Gly His Ser Gly Glu Tyr Gly His Ser Ser Glu Phe Thr Val Asn Phe 305 310 315 320

Lys Glu Lys Leu Thr Gly Lys Leu Phe Ser Asn Leu Gln Asp Arg 325 330 335

His Lys Gly Asn Val Thr Lys Thr Glu Arg Tyr Asp Ile Asp Ala Asn 345 Ile His Gly Asn Arg Phe Arg Gly Ser Ala Thr Ala Ser Asn Lys Asn 360 Asp Thr Ser Lys His Pro Phe Thr Ser Asp Ala Asn Asn Arg Leu Glu Gly Gly Phe Tyr Gly Pro Lys Gly Glu Glu Leu Ala Gly Lys Phe Leu 395 Thr Asn Asp Asn Lys Leu Phe Gly Val Phe Gly Ala Lys Arg Glu Ser 405 410 Lys Ala Glu Glu Lys Thr Glu Ala Ile Leu Asp Ala Tyr Ala Leu Gly 425 Thr Phe Asn Thr Ser Asn Ala Thr Thr Phe Thr Pro Phe Thr Glu Lys 435 440 Gln Leu Asp Asn Phe Gly Asn Ala Lys Lys Leu Val Leu Gly Ser Thr Val Ile Asp Leu Val Pro Thr Asp Ala Thr Lys Asn Glu Phe Thr Lys 475 Asp Lys Pro Glu Ser Ala Thr Asn Glu Ala Gly Glu Thr Leu Met Val 490 Asn Asp Glu Val Ser Val Lys Thr Tyr Gly Lys Asn Phe Glu Tyr Leu Lys Phe Gly Glu Leu Ser Ile Gly Gly Ser His Ser Val Phe Leu Gln 520 Gly Glu Arg Thr Ala Thr Thr Gly Glu Lys Ala Val Pro Thr Thr Gly 530 Thr Ala Lys Tyr Leu Gly Asn Trp Val Gly Tyr Ile Thr Gly Lys Asp Thr Gly Thr Gly Lys Ser Phe Thr Asp Ala Gln Asp Val Ala Asp Phe Asp Ile Asp Phe Gly Asn Lys Ser Val Ser Gly Lys Leu Ile 585 Thr Lys Gly Arg Gln Asp Pro Val Phe Ser Ile Thr Gly Gln Ile Ala 600 Gly Asn Gly Trp Thr Gly Thr Ala Ser Thr Thr Lys Ala Asp Ala Gly Gly Tyr Lys Ile Asp Ser Ser Ser Thr Gly Lys Ser Ile Ala Ile Lys 630 635

Asp Ala Asn Val Thr Gly Gly Phe Tyr Gly Pro Asn Ala Asn Glu Met  $645 \hspace{1cm} 650 \hspace{1cm} 655$ 

Gly Gly Ser Phe Thr His Asn Ala Asp Asp Ser Lys Ala Ser Val Val 660 665 670

Phe Gly Thr Lys Arg Gln Gln Glu Val Lys 675 680

<210> 13

<211> 1070

<212> PRT

<213> Moraxella catarrhalis

<400> 13

Met Asn Gln Ser Lys Lys Ser Lys Ser Lys Gln Val Leu Lys Leu 1 5 10 15

Ser Ala Leu Ser Leu Gly Leu Leu Asn Ile Thr Gln Val Ala Leu Ala 20 25 30

Asn Thr Thr Ala Asp Lys Ala Glu Ala Thr Asp Lys Thr Asn Leu Val
35 40 45

Val Val Leu Asp Glu Thr Val Val Thr Ala Lys Lys Asn Ala Arg Lys
50 60

Ala Asn Glu Val Thr Gly Leu Gly Lys Val Val Lys Thr Ala Glu Thr 65 70 75 80

Ile Asn Lys Glu Gln Val Leu Asn Ile Arg Asp Leu Thr Arg Tyr Asp 85 90 95

Pro Gly Ile Ala Val Val Glu Gln Gly Arg Gly Ala Ser Ser Gly Tyr
100 105 110

Ser Ile Arg Gly Met Asp Lys Asn Arg Val Ala Val Leu Val Asp Gly
115 120 125

Ile Asn Gln Ala Gln His Tyr Ala Leu Gln Gly Pro Val Ala Gly Lys
130 135 140

Asn Tyr Ala Ala Gly Gly Ala Ile Asn Glu Ile Glu Tyr Glu Asn Val 145 150 155 160

Arg Ser Val Glu Ile Ser Lys Gly Ala Asn Ser Ser Glu Tyr Gly Ser 165 170 175

Gly Ala Leu Ser Gly Ser Val Ala Phe Val Thr Lys Thr Ala Asp Asp 180 185 190

Ile Ile Lys Asp Gly Lys Asp Trp Gly Val Gln Thr Lys Thr Ala Tyr
195 200 205

Ala Ser Lys Asn Asn Ala Trp Val Asn Ser Val Ala Ala Ala Gly Lys 210 215 220

Ala Gly Ser Phe Ser Gly Leu Ile Ile Tyr Thr Asp Arg Arg Gly Gln 230 225 235 Glu Tyr Lys Ala His Asp Asp Ala Tyr Gln Gly Ser Gln Ser Phe Asp Arg Ala Val Ala Thr Thr Asp Pro Asn Asn Pro Lys Phe Leu Ile Ala Asn Glu Cys Ala Asn Gly Asn Tyr Glu Ala Cys Ala Ala Gly Gly Gln 280 Thr Lys Leu Gln Ala Lys Pro Thr Asn Val Arg Asp Lys Val Asn Val Lys Asp Tyr Thr Gly Pro Asn Arg Leu Ile Pro Asn Pro Leu Thr Gln 315 Asp Ser Lys Ser Leu Leu Leu Arg Pro Gly Tyr Gln Leu Asn Asp Lys His Tyr Val Gly Gly Val Tyr Glu Ile Thr Lys Gln Asn Tyr Ala Met 345 Gln Asp Lys Thr Val Pro Ala Tyr Leu Thr Val His Asp Ile Glu Lys 355 360 Ser Arg Leu Ser Asn His Gly Gln Ala Asn Gly Tyr Tyr Gln Gly Asn Asn Leu Gly Glu Arg Ile Arg Asp Ala Ile Gly Ala Asn Ser Gly Tyr 385 390 395 Gly Ile Asn Tyr Ala His Gly Val Phe Tyr Asp Glu Lys His Gln Lys Asp Arg Leu Gly Leu Glu Tyr Val Tyr Asp Ser Lys Gly Glu Asn Lys Trp Phe Asp Asp Val Arg Val Ser Tyr Asp Lys Gln Asp Ile Thr Leu Arg Ser Gln Leu Thr Asn Thr His Cys Ser Thr Tyr Pro His Ile Asp 455 Lys Asn Cys Thr Pro Asp Val Asn Lys Pro Phe Ser Val Lys Glu Val Asp Asn Asn Ala Tyr Lys Glu Gln His Asn Leu Ile Lys Ala Val Phe 490 Asn Lys Lys Met Ala Leu Gly Asn Thr His His His Ile Asn Leu Gln 505 Val Gly Tyr Asp Lys Phe Asn Ser Ser Leu Ser Arg Glu Asp Tyr Arg 520 Leu Ala Thr His Gln Ser Tyr Gln Lys Leu Asp Tyr Thr Pro Pro Ser

535

530

Asn Pro Leu Pro Asp Lys Phe Lys Pro Ile Leu Gly Ser Asn Asn Arq 555 Pro Ile Cys Leu Asp Ala Tyr Gly Tyr Gly His Asp His Pro Gln Ala Cys Asn Ala Lys Asn Ser Thr Tyr Gln Asn Phe Ala Ile Lys Lys Gly 585 Ile Glu Gln Tyr Asn Gln Thr Asn Thr Asp Lys Ile Asp Tyr Gln Ala 600 Val Ile Asp Gln Tyr Asp Lys Gln Asn Pro Asn Ser Thr Leu Lys Pro 615 Phe Glu Lys Ile Lys Gln Ser Leu Gly Gln Glu Lys Tyr Asp Glu Ile Asp Arg Leu Gly Phe Asn Ala Tyr Lys Asp Leu Arg Asn Glu Trp Ala Gly Trp Thr Asn Asp Asn Ser Gln Gln Asn Ala Asn Lys Gly Thr Asp 665 Asn Ile Tyr Gln Pro Asn Gln Ala Thr Val Val Lys Asp Asp Lys Cys Lys Tyr Ser Glu Thr Asn Ser Tyr Ala Asp Cys Ser Thr Thr Arg His 695 Ile Ser Gly Asp Asn Tyr Phe Ile Ala Leu Lys Asp Asn Met Thr Ile Asn Lys Tyr Val Asp Leu Gly Leu Gly Ala Arg Tyr Asp Arg Ile Lys His Lys Ser Asp Val Pro Leu Val Asp Asn Ser Ala Ser Asn Gln Leu 740 Ser Trp Asn Phe Gly Val Val Lys Pro Thr Asn Trp Leu Asp Ile Ala Tyr Arg Ser Ser Gln Gly Phe Arg Met Pro Ser Phe Ser Glu Met 770 Tyr Gly Glu Arg Phe Gly Val Thr Ile Gly Lys Gly Thr Gln His Gly 790 Cys Lys Gly Leu Tyr Tyr Ile Cys Gln Gln Thr Val His Gln Thr Lys 805 Leu Lys Pro Glu Lys Ser Phe Asn Gln Glu Ile Gly Ala Thr Leu His Asn His Leu Gly Ser Leu Glu Val Ser Tyr Phe Lys Asn Arg Tyr Thr 840

Asp Leu Ile Val Gly Lys Ser Glu Glu Ile Arg Thr Leu Thr Gln Gly 850 855

Asp Asn Ala Gly Lys Gln Arg Gly Lys Gly Asp Leu Gly Phe His Asn 865 870 875 880

Gly Gln Asp Ala Asp Leu Thr Gly Ile Asn Ile Leu Gly Arg Leu Asp 885 890 895

Leu Asn Ala Val Asn Ser Arg Leu Pro Tyr Gly Leu Tyr Ser Thr Leu 900 905 910

Ala Tyr Asn Lys Val Asp Val Lys Gly Lys Thr Leu Asn Pro Thr Leu 915 920 925

Ala Gly Thr Asn Ile Leu Phe Asp Ala Ile Gln Pro Ser Arg Tyr Val 930 935 940

Val Gly Leu Gly Tyr Asp Ala Pro Ser Gln Lys Trp Gly Ala Asn Ala 945 950 955 960

Ile Phe Thr His Ser Asp Ala Lys Asn Pro Ser Glu Leu Leu Ala Asp 965 970 975

Lys Asn Leu Gly Asn Gly Asn Gln Thr Lys Gln Ala Thr Lys Ala 980 985 990

Lys Ser Thr Pro Trp Gln Thr Leu Asp Leu Ser Gly Tyr Val Asn Ile 995 1000 1005

Lys Asp Asn Phe Thr Leu Arg Ala Gly Val Tyr Asn Val Phe Asn Thr 1010 1015 1020

Tyr Tyr Thr Trp Glu Ala Leu Arg Gln Thr Ala Glu Gly Ala Val 1025 1030 1035 1040

Asn Gln His Thr Gly Leu Ser Gln Asp Lys His Tyr Gly Arg Tyr Ala 1045 1050 1055

Ala Pro Gly Arg Asn Tyr Gln Leu Ala Leu Glu Met Lys Phe 1060 1065 1070

<210> 14

<211> 17

<212> PRT

<213> Moraxella catarrhalis

<400> 14

Asn Gln Ser Lys Lys Ser Lys Ser Lys Gln Val Leu Lys Leu Ser 1 5 10 15

Ala

<210> 15

<211> 709

<212> PRT

<213> Moraxella catarrhalis

<400> 15

Met Lys His Ile Pro Leu Thr Thr Leu Cys Val Ala Ile Ser Ala Val 1 5 10 15

Leu Leu Thr Ala Cys Gly Gly Ser Ser Gly Gly Phe Asn Pro Pro Ala
20 25 30

Ser Thr Pro Ile Pro Asn Ala Gly Asn Ser Gly Asn Ala Gly Asn Ala 35 40 45

Gly Asn Ala Gly Gly Thr Gly Gly Ala Asn Ser Gly Ala Gly Asn Ala
50 55 60

Gly Gly Thr Gly Gly Ala Asn Ser Gly Ala Gly Ser Ala Ser Thr Pro 65 70 75 80

Glu Pro Lys Tyr Lys Asp Val Pro Thr Asp Glu Asn Lys Lys Ala Glu 85 90 95

Val Ser Gly Ile Gln Glu Pro Ala Met Gly Tyr Gly Val Glu Leu Lys 100 105 110

Leu Arg Asn Trp Ile Pro Gln Glu Gln Glu Glu His Ala Lys Ile Asn 115 120 125

Thr Asn Asp Val Val Lys Leu Glu Gly Asp Leu Lys His Asn Pro Phe 130 135 140

Asp Asn Ser Ile Trp Gln Asn Ile Lys Asn Ser Lys Glu Val Gln Thr 145 150 155 160

Val Tyr Asn Gln Glu Lys Gln Asn Ile Glu Asp Gln Ile Lys Arg Glu 165 170 175

Asn Lys Gln Arg Pro Asp Lys Lys Leu Asp Asp Val Ala Leu Gln Ala 180 185 190

Tyr Ile Glu Lys Val Leu Asp Asp Arg Leu Thr Glu Leu Ala Lys Pro 195 200 205

Ile Tyr Glu Lys Asn Ile Asn Tyr Ser His Asp Lys Gln Asn Lys Ala 210 215 220

Arg Thr Arg Asp Leu Lys Tyr Val Arg Ser Gly Tyr Ile Tyr Arg Ser 225 230 235 240

Gly Tyr Ser Asn Ile Ile Pro Lys Lys Ile Ala Lys Thr Gly Phe Asp 245 250 255

Gly Ala Leu Phe Tyr Gln Gly Thr Gln Thr Ala Lys Gln Leu Pro Val 260 265 270

Ser Gln Val Lys Tyr Lys Gly Thr Trp Asp Phe Met Thr Asp Ala Lys
275
280
285

Lys Gly Gln Ser Phe Ser Ser Phe Gly Thr Ser Gln Arg Leu Ala Gly 290 295 300

Asp Arg Tyr Ser Ala Met Ser Tyr His Glu Tyr Pro Ser Leu Leu Thr 305 310 315 320

Asp Glu Lys Asn Lys Pro Asp Asn Tyr Asn Gly Glu Tyr Gly His Ser 325 330 335

Ser Glu Phe Thr Val Asp Phe Ser Lys Lys Ser Leu Lys Gly Glu Leu 340 345 350

Ser Ser Asn Ile Gln Asp Gly His Lys Gly Ser Val Asn Lys Thr Lys 355 360 365

Arg Tyr Asp Ile Asp Ala Asn Ile Tyr Gly Asn Arg Phe Arg Gly Ser 370 380

Ala Thr Ala Ser Asp Thr Thr Glu Ala Ser Lys Ser Lys His Pro Phe 385 390 395 400

Thr Ser Asp Ala Lys Asn Ser Leu Glu Gly Gly Phe Tyr Gly Pro Asn 405 410 415

Ala Glu Glu Leu Ala Gly Lys Phe Leu Thr Asn Asp Asn Lys Leu Phe
420 425 430

Gly Val Phe Gly Ala Lys Arg Glu Ser Glu Ala Lys Glu Lys Thr Glu 435 440 445

Ala Ile Leu Asp Ala Tyr Ala Leu Gly Thr Phe Asn Lys Pro Gly Thr 450 455 460

Thr Asn Pro Ala Phe Thr Ala Asn Ser Lys Lys Glu Leu Asp Asn Phe 465 470 475 480

Gly Asn Ala Lys Lys Leu Val Leu Gly Ser Thr Val Ile Asp Leu Val
485 490 495

Pro Thr Gly Ala Thr Lys Asp Val Asn Glu Phe Lys Glu Lys Pro Lys 500 505 510

Ser Ala Thr Asn Lys Ala Gly Glu Thr Leu Met Val Asn Asp Glu Val 515 520 525

Ile Val Lys Thr Tyr Gly Tyr Gly Arg Asn Phe Glu Tyr Leu Lys Phe 530 535 540

Gly Glu Leu Ser Ile Gly Gly Ser His Ser Val Phe Leu Gln Gly Glu 545 550 555

Arg Thr Ala Glu Lys Ala Val Pro Thr Glu Gly Thr Ala Lys Tyr Leu 565 570 575

Gly Asn Trp Val Gly Tyr Ile Thr Gly Lys Asp Thr Gly Thr Ser Thr 580 585 590

Gly Lys Ser Phe Asn Glu Ala Gln Asp Ile Ala Asp Phe Asp Ile Asp 595 600 605

Phe Glu Arg Lys Ser Val Lys Gly Lys Leu Thr Thr Gln Gly Arg Gln 610 620

Asp Pro Val Phe Asn Ile Thr Gly Gln Ile Ala Gly Asn Gly Trp Thr 625 630 635 640

Gly Thr Ala Ser Thr Ala Lys Ala Asn Val Gly Gly Tyr Lys Ile Asp 645 650 655

Ser Ser Ser Thr Gly Lys Ser Ile Val Ile Glu Asn Ala Lys Val Thr 660 665 670

Gly Gly Phe Tyr Gly Pro Asn Ala Asn Glu Met Gly Gly Ser Phe Thr 675 680 685

His Asp Thr Asp Asp Ser Lys Ala Ser Val Val Phe Gly Thr Lys Arg 690 695 700

Gln Glu Glu Val Lys 705

<210> 16

<211> 689

<212> PRT

<213> Moraxella catarrhalis

<400> 16

Cys Gly Gly Ser Ser Gly Gly Phe Asn Pro Pro Ala Ser Thr Pro Ile 1 5 10 15

Pro Asn Ala Gly Asn Ser Gly Asn Ala Gly Asn Ala Gly 20 25 30

Gly Thr Gly Gly Ala Asn Ser Gly Ala Gly Asn Ala Gly Gly Thr Gly 35 40 45

Gly Ala Asn Ser Gly Ala Gly Ser Ala Ser Thr Pro Glu Pro Lys Tyr 50 55 60

Lys Asp Val Pro Thr Asp Glu Asn Lys Lys Ala Glu Val Ser Gly Ile 65 70 75 80

Gln Glu Pro Ala Met Gly Tyr Gly Val Glu Leu Lys Leu Arg Asn Trp 85 90 95

Ile Pro Gln Glu Glu Glu His Ala Lys Ile Asn Thr Asn Asp Val 100 105 110

Val Lys Leu Glu Gly Asp Leu Lys His Asn Pro Phe Asp Asn Ser Ile 115 120 125

Trp Gln Asn Ile Lys Asn Ser Lys Glu Val Gln Thr Val Tyr Asn Gln 130 135 140

Glu Lys Gln Asn Ile Glu Asp Gln Ile Lys Arg Glu Asn Lys Gln Arg 145 150 155 160

Pro Asp Lys Lys Leu Asp Asp Val Ala Leu Gln Ala Tyr Ile Glu Lys 170 Val Leu Asp Asp Arg Leu Thr Glu Leu Ala Lys Pro Ile Tyr Glu Lys 185 Asn Ile Asn Tyr Ser His Asp Lys Gln Asn Lys Ala Arg Thr Arg Asp Leu Lys Tyr Val Arg Ser Gly Tyr Ile Tyr Arg Ser Gly Tyr Ser Asn Ile Ile Pro Lys Lys Ile Ala Lys Thr Gly Phe Asp Gly Ala Leu Phe Tyr Gln Gly Thr Gln Thr Ala Lys Gln Leu Pro Val Ser Gln Val Lys 250 Tyr Lys Gly Thr Trp Asp Phe Met Thr Asp Ala Lys Lys Gly Gln Ser 260 265 Phe Ser Ser Phe Gly Thr Ser Gln Arg Leu Ala Gly Asp Arg Tyr Ser Ala Met Ser Tyr His Glu Tyr Pro Ser Leu Leu Thr Asp Glu Lys Asn 290 Lys Pro Asp Asn Tyr Asn Gly Glu Tyr Gly His Ser Ser Glu Phe Thr Val Asp Phe Ser Lys Lys Ser Leu Lys Gly Glu Leu Ser Ser Asn Ile 330 Gln Asp Gly His Lys Gly Ser Val Asn Lys Thr Lys Arg Tyr Asp Ile Asp Ala Asn Ile Tyr Gly Asn Arg Phe Arg Gly Ser Ala Thr Ala Ser Asp Thr Thr Glu Ala Ser Lys Ser Lys His Pro Phe Thr Ser Asp Ala Lys Asn Ser Leu Glu Gly Gly Phe Tyr Gly Pro Asn Ala Glu Glu Leu Ala Gly Lys Phe Leu Thr Asn Asp Asn Lys Leu Phe Gly Val Phe Gly 405 Ala Lys Arg Glu Ser Glu Ala Lys Glu Lys Thr Glu Ala Ile Leu Asp 425 Ala Tyr Ala Leu Gly Thr Phe Asn Lys Pro Gly Thr Thr Asn Pro Ala 435 440 Phe Thr Ala Asn Ser Lys Lys Glu Leu Asp Asn Phe Gly Asn Ala Lys Lys Leu Val Leu Gly Ser Thr Val Ile Asp Leu Val Pro Thr Gly Ala 475

Thr Lys Asp Val Asn Glu Phe Lys Glu Lys Pro Lys Ser Ala Thr Asn 485 490 495

Lys Ala Gly Glu Thr Leu Met Val Asn Asp Glu Val Ile Val Lys Thr
500 505 510

Tyr Gly Tyr Gly Arg Asn Phe Glu Tyr Leu Lys Phe Gly Glu Leu Ser 515 520 525

Ile Gly Gly Ser His Ser Val Phe Leu Gln Gly Glu Arg Thr Ala Glu 530 540

Lys Ala Val Pro Thr Glu Gly Thr Ala Lys Tyr Leu Gly Asn Trp Val 545 550 555 560

Gly Tyr Ile Thr Gly Lys Asp Thr Gly Thr Ser Thr Gly Lys Ser Phe 565 570 575

Asn Glu Ala Gln Asp Ile Ala Asp Phe Asp Ile Asp Phe Glu Arg Lys 580 585 590

Ser Val Lys Gly Lys Leu Thr Thr Gln Gly Arg Gln Asp Pro Val Phe 595 600 605

Asn Ile Thr Gly Gln Ile Ala Gly Asn Gly Trp Thr Gly Thr Ala Ser 610 620

Thr Ala Lys Ala Asn Val Gly Gly Tyr Lys Ile Asp Ser Ser Thr 625 630 635

Gly Lys Ser Ile Val Ile Glu Asn Ala Lys Val Thr Gly Gly Phe Tyr \$645\$ \$650\$

Gly Pro Asn Ala Asn Glu Met Gly Gly Ser Phe Thr His Asp Thr Asp 660 665 670

Asp Ser Lys Ala Ser Val Val Phe Gly Thr Lys Arg Gln Glu Glu Val 675 680 685

Lys

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<213> Moraxella catarrhalis

<400> 17

Asn Glu Val Thr Gly Leu Gly

<210> 18

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<400> 20
aaacacattc ctttaaccac actgtgtgtg gcaatctctg ccgtcttatt aaccgct
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<212> PRT
<213> Moraxella catarrhalis
<400> 21
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Ile Gly Cys Tyr Val Lys Ala Glu Thr Gln Ser Ile Lys Asp Thr Lys
Glu Ala Ile Ser Ser Glu Val Asp Thr Gln Ser Thr Glu Asp Ser Glu
Leu Glu Thr Ile Ser Val Thr Ala Glu Lys Ile Arg Asp Arg Lys Asp
Asn Glu Val Thr Gly Leu Gly Lys Ile Ile Lys Thr Ser Glu Ser Ile
Ser Arg Glu Gln Val Leu Asn Ile Arg Asp Leu Thr Arg Tyr Asp Pro
Gly Ile Ser Val Val Glu Gln Gly Arg Gly Ala Ser Ser Gly Tyr Ser
            100
                                105
Ile Arg Gly Met Asp Arg Asn Arg Val Ala Leu Leu Val Asp Gly Leu
                            120
Pro Gln Thr Gln Ser Tyr Val Val Gln Ser Pro Leu Val Ala Arg Ser
    130
                        135
Gly Tyr Ser Gly Thr Gly Ala Ile Asn Glu Ile Glu Tyr Glu Asn Val
                    150
                                         155
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Lys Ala Val Glu Ile Ser Lys Gly Gly Ser Ser Ser Glu Tyr Gly Asn Gly Ala Leu Ala Gly Ser Val Thr Phe Gln Ser Lys Ser Ala Ala Asp 185 Ile Leu Glu Gly Asp Lys Ser Trp Gly Ile Gln Thr Lys Asn Ala Tyr 200 Ser Ser Lys Asn Lys Gly Phe Thr His Ser Leu Ala Val Ala Gly Lys Gln Gly Gly Phe Glu Gly Leu Ala Ile Tyr Thr Gln Arg Asn Ser Ile Glu Thr Gln Val His Lys Asp Ala Leu Lys Gly Val Gln Ser Tyr Asp 245 250 Arg Leu Ile Ala Thr Thr Asp Lys Ser Ser Gly Tyr Phe Val Ile Gln Gly Glu Cys Pro Asn Gly Asp Asp Lys Cys Ala Ala Lys Pro Pro Ala Thr Leu Ser Thr Gln Ser Glu Thr Val Ser Val Ser Asp Tyr Thr Gly 295 Ala Asn Arg Ile Lys Pro Asn Pro Met Lys Tyr Glu Ser Gln Ser Trp 310 315 Phe Leu Arg Gly Gly Tyr His Phe Ser Glu Gln His Tyr Ile Gly Gly 325 Ile Phe Glu Phe Thr Gln Gln Lys Phe Asp Ile Arg Asp Met Thr Phe 345 Pro Ala Tyr Leu Ser Pro Thr Glu Arg Arg Asp Asp Ser Ser Arg Ser 355 Phe Tyr Pro Met Gln Asp His Gly Ala Tyr Gln His Ile Glu Asp Gly Arg Gly Val Lys Tyr Ala Ser Gly Leu Tyr Phe Asp Glu His His Arg 395 Lys Gln Arg Val Gly Ile Glu Tyr Ile Tyr Glu Asn Lys Asn Lys Ala Gly Ile Ile Asp Lys Ala Val Leu Ser Ala Asn Gln Gln Asn Ile Ile 425 Leu Asp Ser Tyr Met Arg His Thr His Cys Ser Leu Tyr Pro Asn Pro Ser Lys Asn Cys Arg Pro Thr Leu Asp Lys Pro Tyr Ser Tyr Tyr Arg 455 Ser Asp Arg Asn Val Tyr Lys Glu Lys His Asn Met Leu Gln Leu Asn 470 475

Leu Glu Lys Lys Ile Gln Gln Asn Trp Leu Thr His Gln Ile Val Phe 485 490 495

Asn Leu Gly Phe Asp Asp Phe Thr Ser Ala Leu Gln His Lys Asp Tyr 500 505 510

Leu Thr Arg Arg Val Ile Ala Thr Ala Asp Ser Ile Pro Arg Lys Pro 515 520 525

Gly Glu Thr Gly Lys Pro Arg Asn Gly Leu Gln Ser Gln Pro Tyr Leu 530 540

Tyr Pro Lys Pro Glu Pro Tyr Phe Ala Gly Gln Asp His Cys Asn Tyr 545 550 555 560

Gln Gly Ser Ser Ser Asn Tyr Arg Asp Cys Lys Val Arg Leu Ile Lys
565 570 575

Gly Lys Asn Tyr Tyr Phe Ala Ala Arg Asn Asn Met Ala Leu Gly Lys 580 585 590

Tyr Val Asp Leu Gly Leu Gly Ile Arg Tyr Asp Val Ser Arg Thr Lys 595 600 605

Ala Asn Glu Ser Thr Ile Ser Val Gly Lys Phe Lys Asn Phe Ser Trp 610 615 620

Asn Thr Gly Ile Val Ile Lys Pro Thr Glu Trp Leu Asp Leu Ser Tyr 625 630 635 640

Arg Leu Ser Thr Gly Phe Arg Asn Pro Ser Phe Ser Glu Met Tyr Gly 645 650 655

Trp Arg Tyr Gly Gly Lys Asn Asp Glu Val Tyr Val Gly Lys Phe Lys 660 665 670

Pro Glu Thr Ser Arg Asn Gln Glu Phe Gly Leu Ala Leu Lys Gly Asp 675 680 685

Phe Gly Asn Ile Glu Ile Ser His Phe Ser Asn Ala Tyr Arg Asn Leu 690 695 700

Ile Ala Phe Ala Glu Glu Leu Ser Lys Asn Gly Thr Gly Lys Gly Asn 705 710 715 720

Tyr Gly Tyr His Asn Ala Gln Asn Ala Lys Leu Val Gly Val Asn Ile 725 730 735

Thr Ala Gln Leu Asp Phe Asn Gly Leu Trp Lys Arg Ile Pro Tyr Gly
740 745 750

Trp Tyr Ala Thr Phe Ala Tyr Asn Gln Val Lys Val Lys Asp Gln Lys
755 760 765

Ile Asn Ala Gly Leu Ala Ser Val Ser Ser Tyr Leu Phe Asp Ala Ile 770 775 780 Gln Pro Ser Arg Tyr Ile Ile Gly Leu Gly Tyr Asp His Pro Ser Asn 785 790 795 800

Thr Trp Gly Ile Asn Thr Met Phe Thr Gln Ser Lys Ala Lys Ser Gln 805 810 815

Asn Glu Leu Gly Lys Arg Ala Leu Gly Asn Asn Ser Arg Asp Val

Lys Ser Thr Arg Lys Leu Thr Arg Ala Trp His Ile Leu Asp Val Ser 835 840 845

Gly Tyr Tyr Met Ala Asn Lys Asn Ile Met Leu Arg Leu Gly Ile Tyr 850 855 860

Asn Leu Phe Asn Tyr Arg Tyr Val Thr Trp Glu Ala Val Arg Gln Thr 865 870 875 880

Ala Gln Gly Ala Val Asn Gln His Gln Asn Val Gly Ser Tyr Thr Arg 885 890 895

Tyr Ala Ala Ser Gly Arg Asn Tyr Thr Leu Thr Leu Glu Met Lys Phe 900 905 910

<210> 22

<211> 908

<212> PRT

<213> Moraxella catarrhalis

<400> 22

Met Gln Gln His Leu Phe Arg Leu Asn Ile Leu Cys Leu Ser Leu
1 5 10 15

Met Thr Ala Leu Pro Val Tyr Ala Glu Asn Val Gln Ala Glu Gln Ala 20 25 30

Gln Glu Lys Gln Leu Asp Thr Ile Gln Val Lys Ala Lys Lys Gln Lys
35 40 45

Thr Arg Arg Asp Asn Glu Val Thr Gly Leu Gly Lys Leu Val Lys Ser 50 60

Ser Asp Thr Leu Ser Lys Glu Gln Val Leu Asn Ile Arg Asp Leu Thr 65 70 75 80

Arg Tyr Asp Pro Gly Ile Ala Val Val Glu Gln Gly Arg Gly Ala Ser 85 90 95

Ser Gly Tyr Ser Ile Arg Gly Met Asp Lys Asn Arg Val Ser Leu Thr 100 105 110

Val Asp Gly Val Ser Gln Ile Gln Ser Tyr Thr Ala Gln Ala Leu 115 120 125

Gly Gly Thr Arg Thr Ala Gly Ser Ser Gly Ala Ile Asn Glu Ile Glu Tyr Glu Asn Val Lys Ala Val Glu Ile Ser Lys Gly Ser Asn Ser Ser Glu Tyr Gly Asn Gly Ala Leu Ala Gly Ser Val Ala Phe Gln Thr Lys 170 Thr Ala Ala Asp Ile Ile Gly Glu Gly Lys Gln Trp Gly Ile Gln Ser Lys Thr Ala Tyr Ser Gly Lys Asp His Ala Leu Thr Gln Ser Leu Ala Leu Ala Gly Arg Ser Gly Gly Ala Glu Ala Leu Leu Ile Tyr Thr Lys 215 Arg Arg Gly Arg Glu Ile His Ala His Lys Asp Ala Gly Lys Gly Val Gln Ser Phe Asn Arg Leu Val Leu Asp Glu Asp Lys Lys Glu Gly Gly Ser Gln Tyr Arg Tyr Phe Ile Val Glu Glu Glu Cys His Asn Gly Tyr Ala Ala Cys Lys Asn Lys Leu Lys Glu Asp Ala Ser Val Lys Asp Glu Arg Lys Thr Val Ser Thr Gln Asp Tyr Thr Gly Ser Asn Arg Leu Leu 295 Ala Asn Pro Leu Glu Tyr Gly Ser Gln Ser Trp Leu Phe Arg Pro Gly 315 Trp His Leu Asp Asn Arg His Tyr Val Gly Ala Val Leu Glu Arg Thr 325 Gln Gln Thr Phe Asp Thr Arg Asp Met Thr Val Pro Ala Tyr Phe Thr Ser Glu Asp Tyr Val Pro Gly Ser Leu Lys Gly Leu Gly Lys Tyr Ser Gly Asp Asn Lys Ala Glu Arg Leu Phe Val Gln Gly Glu Gly Ser Thr Leu Gln Gly Ile Gly Tyr Gly Thr Gly Val Phe Tyr Asp Glu Arg His 395 Thr Lys Asn Arg Tyr Gly Val Glu Tyr Val Tyr His Asn Ala Asp Lys 405 Asp Thr Trp Ala Asp Tyr Ala Arg Leu Ser Tyr Asp Arg Gln Gly Ile 425 Asp Leu Asp Asn Arg Leu Gln Gln Thr His Cys Ser His Asp Gly Ser 435 440

Asp Lys Asn Cys Arg Pro Asp Gly Asn Lys Pro Tyr Ser Phe Tyr Lys 450 Ser Asp Arg Met Ile Tyr Glu Glu Ser Arg Asn Leu Phe Gln Ala Val 475 Phe Lys Lys Ala Phe Asp Thr Ala Lys Ile Arg His Asn Leu Ser Ile 490 Asn Leu Gly Tyr Asp Arg Phe Lys Ser Gln Leu Ser His Ser Asp Tyr 500 Tyr Leu Gln Asn Ala Val Gln Ala Tyr Asp Leu Ile Thr Pro Pro Lys 520 Pro Pro Phe Pro Asn Gly Ser Lys Asp Asn Pro Tyr Arg Val Ser Ile Gly Lys Thr Thr Val Asn Thr Ser Pro Ile Cys Arg Phe Gly Asn Asn 550 Thr Tyr Thr Asp Cys Thr Pro Arg Asn Ile Gly Gly Asn Gly Tyr Tyr Ala Ala Val Gln Asp Asn Val Arg Leu Gly Arg Trp Ala Asp Val Gly Ala Gly Ile Arg Tyr Asp Tyr Arg Ser Thr His Ser Glu Asp Lys Ser Val Ser Thr Gly Thr His Arg Asn Leu Ser Trp Asn Ala Gly Val Val Leu Lys Pro Phe Thr Trp Met Asp Leu Thr Tyr Arg Ala Ser Thr Gly 630 Phe Arg Leu Pro Ser Phe Ala Glu Met Tyr Gly Trp Arg Ala Gly Glu Ser Leu Lys Thr Leu Asp Leu Lys Pro Glu Lys Ser Phe Asn Arg Glu Ala Gly Ile Val Phe Lys Gly Asp Phe Gly Asn Leu Glu Ala Ser Tyr Phe Asn Asn Ala Tyr Arg Asp Leu Ile Ala Phe Gly Tyr Glu Thr Arg 690 Thr Gln Asn Gly Gln Thr Ser Ala Ser Gly Asp Pro Gly Tyr Arg Asn Ala Gln Asn Ala Arg Ile Ala Gly Ile Asn Ile Leu Gly Lys Ile Asp 730 Trp His Gly Val Trp Gly Gly Leu Pro Asp Gly Leu Tyr Ser Thr Leu

Ala Tyr Asn Arg Ile Lys Val Lys Asp Ala Asp Ile Arg Ala Asp Arg 755 760 765

Thr Phe Val Thr Ser Tyr Leu Phe Asp Ala Val Gln Pro Ser Arg Tyr 770 775 780

Val Leu Gly Leu Gly Tyr Asp His Pro Asp Gly Ile Trp Gly Ile Asn 785 790 795 800

Thr Met Phe Thr Tyr Ser Lys Ala Lys Ser Val Asp Glu Leu Leu Gly 805 810 815

Ser Gln Ala Leu Leu Asn Gly Asn Ala Asn Ala Lys Lys Ala Ala Ser 820 825 830

Arg Arg Thr Arg Pro Trp Tyr Val Thr Asp Val Ser Gly Tyr Tyr Asn 835 840 845

Ile Lys Lys His Leu Thr Leu Arg Ala Gly Val Tyr Asn Leu Leu Asn 850 855 860

Tyr Arg Tyr Val Thr Trp Glu Asn Val Arg Gln Thr Ala Gly Gly Ala 865 870 875 880

Val Asn Gln His Lys Asn Val Gly Val Tyr Asn Arg Tyr Ala Ala Pro 885 890 895

Gly Arg Asn Tyr Thr Phe Ser Leu Glu Met Lys Phe 900 905

<210> 23

<211> 911

<212> PRT

<213> Moraxella catarrhalis

<400> 23

Met Gln Gln His Leu Phe Arg Leu Asn Ile Leu Cys Leu Ser Leu 1 5 10 15

Met Thr Ala Leu Pro Ala Tyr Ala Glu Asn Val Gln Ala Gly Gln Ala 20 25 30

Gln Glu Lys Gln Leu Asp Thr Ile Gln Val Lys Ala Lys Lys Gln Lys 35 40 45

Thr Arg Arg Asp Asn Glu Val Thr Gly Leu Gly Lys Leu Val Lys Thr 50 55 60

Ala Asp Thr Leu Ser Lys Glu Gln Val Leu Asp Ile Arg Asp Leu Thr 65 70 75 80

Arg Tyr Asp Pro Gly Ile Ala Val Val Glu Gln Gly Arg Gly Ala Ser 85 90 95

Ser Gly Tyr Ser Ile Arg Gly Met Asp Lys Asn Arg Val Ser Leu Thr 100 105 110 Val Asp Gly Leu Ala Gln Ile Gln Ser Tyr Thr Ala Gln Ala Ala Leu 115 120 125

Gly Gly Thr Arg Thr Ala Gly Ser Ser Gly Ala Ile Asn Glu Ile Glu 130 135 140

Tyr Glu Asn Val Lys Ala Val Glu Ile Ser Lys Gly Ser Asn Ser Val 145 150 155 160

Glu Gln Gly Ser Gly Ala Leu Ala Gly Ser Val Ala Phe Gln Thr Lys 165 170 175

Thr Ala Asp Asp Val Ile Gly Glu Gly Arg Gln Trp Gly Ile Gln Ser 180 185 190

Lys Thr Ala Tyr Ser Gly Lys Asn Arg Gly Leu Thr Gln Ser Ile Ala 195 200 205

Leu Ala Gly Arg Ile Gly Gly Ala Glu Ala Leu Leu Ile His Thr Gly 210 215 220

Arg Arg Ala Gly Glu Ile Arg Ala His Glu Asp Ala Gly Arg Gly Val 225 230 235 240

Gln Ser Phe Asn Arg Leu Val Pro Val Glu Asp Ser Ser Glu Tyr Ala 245 250 255

Tyr Phe Ile Val Glu Asp Glu Cys Glu Gly Lys Asn Tyr Glu Thr Cys 260 265 270

Lys Ser Lys Pro Lys Lys Asp Val Val Gly Lys Asp Glu Arg Gln Thr 275 280 285

Val Ser Thr Arg Asp Tyr Thr Gly Pro Asn Arg Phe Leu Ala Asp Pro 290 295 300

Leu Ser Tyr Glu Ser Arg Ser Trp Leu Phe Arg Pro Gly Phe Arg Phe 305 310 315 320

Glu Asn Lys Arg His Tyr Ile Gly Gly Ile Leu Glu His Thr Gln Gln 325 330 335

Thr Phe Asp Thr Arg Asp Met Thr Val Pro Ala Phe Leu Thr Lys Ala 340 345 350

Val Phe Asp Ala Asn Ser Lys Gln Ala Gly Ser Leu Pro Gly Asn Gly 355 360 365

Lys Tyr Ala Gly Asn His Lys Tyr Gly Gly Leu Phe Thr Asn Gly Glu 370 375 380

Asn Gly Ala Leu Val Gly Ala Glu Tyr Gly Thr Gly Val Phe Tyr Asp 385 390 395 400

Glu Thr His Thr Lys Ser Arg Tyr Gly Leu Glu Tyr Val Tyr Thr Asn 405 410 415

Ala Asp Lys Asp Thr Trp Ala Asp Tyr Ala Arg Leu Ser Tyr Asp Arg 420 425 430

Gln Gly Ile Gly Leu Asp Asn His Phe Gln Gln Thr His Cys Ser Ala 435 440 445

Asp Gly Ser Asp Lys Tyr Cys Arg Pro Ser Ala Asp Lys Pro Phe Ser 450 460

Tyr Tyr Lys Ser Asp Arg Val Ile Tyr Gly Glu Ser His Arg Leu Leu 465 470 475 480

Gln Ala Ala Phe Lys Lys Ser Phe Asp Thr Ala Lys Ile Arg His Asn 485 490 495

Leu Ser Val Asn Leu Gly Phe Asp Arg Phe Asp Ser Asn Leu Arg His 500 505 510

Gln Asp Tyr Tyr Gln His Ala Asn Arg Ala Tyr Ser Ser Lys Thr 515 520 525

Pro Pro Lys Thr Ala Asn Pro Asn Gly Asp Lys Ser Lys Pro Tyr Trp 530 540

Val Ser Ile Gly Gly Gly Asn Val Val Thr Gly Gln Ile Cys Leu Phe 545 550 555 560

Gly Asn Asn Thr Tyr Thr Asp Cys Thr Pro Arg Ser Ile Asn Gly Lys 565 570 575

Ser Tyr Tyr Ala Ala Val Arg Asp Asn Val Arg Leu Gly Arg Trp Ala 580 585 590

Asp Val Gly Ala Gly Leu Arg Tyr Asp Tyr Arg Ser Thr His Ser Asp 595 600 605

Asp Gly Ser Val Ser Thr Gly Thr His Arg Thr Leu Ser Trp Asn Ala 610 620

Gly Ile Val Leu Lys Pro Ala Asp Trp Leu Asp Leu Thr Tyr Arg Thr 625 630 635 640

Ser Thr Gly Phe Arg Leu Pro Ser Phe Ala Glu Met Tyr Gly Trp Arg 645 650 655

Ser Gly Val Gln Ser Lys Ala Val Lys Ile Asp Pro Glu Lys Ser Phe 660 665 670

Asn Lys Glu Ala Gly Ile Val Phe Lys Gly Asp Phe Gly Asn Leu Glu 675 680 685

Ala Ser Trp Phe Asn Asn Ala Tyr Arg Asp Leu Ile Val Arg Gly Tyr 690 695 700

Glu Ala Gln Ile Lys Asn Gly Lys Glu Glu Ala Lys Gly Asp Pro Ala 705 710 715 720

Tyr Leu Asn Ala Gln Ser Ala Arg Ile Thr Gly Ile Asn Ile Leu Gly
725 730 735

Lys Ile Asp Trp Asn Gly Val Trp Asp Lys Leu Pro Glu Gly Trp Tyr 740 745 750

Ser Thr Phe Ala Tyr Asn Arg Val His Val Arg Asp Ile Lys Lys Arg 755 760 765

Ala Asp Arg Thr Asp Ile Gln Ser His Leu Phe Asp Ala Ile Gln Pro 770 780

Ser Arg Tyr Val Val Gly Leu Gly Tyr Asp Gln Pro Glu Gly Lys Trp 785 790 795 800

Gly Val Asn Gly Met Leu Thr Tyr Ser Lys Ala Lys Glu Ile Thr Glu 805 810 815

Leu Leu Gly Ser Arg Ala Leu Leu Asn Gly Asn Ser Arg Asn Thr Lys
820 825 830

Ala Thr Ala Arg Arg Thr Arg Pro Trp Tyr Ile Val Asp Val Ser Gly 835 840 845

Tyr Tyr Thr Ile Lys Lys His Phe Thr Leu Arg Ala Gly Val Tyr Asn 850 . 855 860

Leu Leu Asn Tyr Arg Tyr Val Thr Trp Glu Asn Val Arg Gln Thr Ala 865 870 875 880

Gly Gly Ala Val Asn Gln His Lys Asn Val Gly Val Tyr Asn Arg Tyr 885 890 895

Ala Ala Pro Gly Arg Asn Tyr Thr Phe Ser Leu Glu Met Lys Phe 900 905 910

<210> 24

<211> 915

<212> PRT

<213> Moraxella catarrhalis

<400> 24

Met Gln Gln His Leu Phe Arg Leu Asn Ile Leu Cys Leu Ser Leu 1 5 10 15

Met Thr Ala Leu Pro Ala Tyr Ala Glu Asn Val Gln Ala Gly Gln Ala 20 25 30

Gln Glu Lys Gln Leu Asp Thr Ile Gln Val Lys Ala Lys Lys Gln Lys 35 40 45

Thr Arg Arg Asp Asn Glu Val Thr Gly Leu Gly Lys Leu Val Lys Thr 50 55 60

Ala Asp Thr Leu Ser Lys Glu Gln Val Leu Asp Ile Arg Asp Leu Thr
65 70 75 80

Arg Tyr Asp Pro Gly Ile Ala Val Val Glu Gln Gly Arg Gly Ala Ser 85 90 95 Ser Gly Tyr Ser Ile Arg Gly Met Asp Lys Asn Arg Val Ser Leu Thr

Val Asp Gly Leu Ala Gln Ile Gln Ser Tyr Thr Ala Gln Ala Leu 115 120 125

Gly Gly Thr Arg Thr Ala Gly Ser Ser Gly Ala Ile Asn Glu Ile Glu 130 135 140

Tyr Glu Asn Val Lys Ala Val Glu Ile Ser Lys Gly Ser Asn Ser Val 145 150 155 160

Glu Gln Gly Ser Gly Ala Leu Ala Gly Ser Val Ala Phe Gln Thr Lys 165 170 175

Thr Ala Asp Asp Val Ile Gly Glu Gly Arg Gln Trp Gly Ile Gln Ser 180 185 190

Lys Thr Ala Tyr Ser Gly Lys Asn Arg Gly Leu Thr Gln Ser Leu Ala 195 200 205

Leu Ala Gly Arg Ile Gly Gly Ala Glu Ala Leu Leu Ile Arg Thr Gly 210 215 220

Arg His Ala Gly Glu Ile Arg Ala His Glu Ala Ala Gly Arg Gly Val 225 230 235 240

Gln Ser Phe Asn Arg Leu Ala Pro Val Asp Asp Gly Ser Lys Tyr Ala 245 250 255

Tyr Phe Ile Val Glu Glu Glu Cys Lys Asn Gly Gly His Glu Lys Cys  $\phantom{0}260$  .  $\phantom{0}265$   $\phantom{0}270$ 

Lys Ala Asn Pro Pro Lys Asp Val Val Gly Glu Asp Lys Arg Gln Thr 275 280 285

Val Ser Thr Arg Asp Tyr Thr Gly Pro Asn Arg Phe Leu Ala Asp Pro 290 295 300

Leu Ser Tyr Glu Ser Arg Ser Trp Leu Phe Arg Pro Gly Phe Arg Phe 305 310 315 320

Glu Asn Lys Arg His Tyr Ile Gly Gly Ile Leu Glu Arg Thr Gln Gln 325 330 335

Thr Phe Asp Thr Arg Asp Met Thr Val Pro Ala Phe Leu Thr Lys Ala 340 345 350

Val Phe Asp Ala Asn Gln Lys Gln Ala Gly Ser Leu Arg Gly Asn Gly 355 360 365

Lys Tyr Ala Gly Asn His Lys Tyr Gly Gly Leu Phe Thr Ser Gly Glu 370 375 380

Asn Asn Ala Pro Val Gly Ala Glu Tyr Gly Thr Gly Val Phe Tyr Asp 385 . 390 395 400

Glu Thr His Thr Lys Ser Arg Tyr Gly Leu Glu Tyr Val Tyr Thr Asn 405 410 415

Ala Asp Lys Asp Thr Trp Ala Asp Tyr Ala Arg Leu Ser Tyr Asp Arg 425 Gln Gly Ile Gly Leu Asp Asn His Phe Gln Gln Thr His Cys Ser Ala Asp Gly Ser Asp Lys Tyr Cys Arg Pro Ser Ala Asp Lys Pro Phe Ser Tyr Tyr Lys Ser Asp Arg Val Ile Tyr Gly Glu Ser His Lys Leu Leu 470 475 Gln Ala Ala Phe Lys Lys Ser Phe Asp Thr Ala Lys Ile Arg His Asn Leu Ser Val Asn Leu Gly Tyr Asp Arg Phe Gly Ser Asn Leu Arg His 500 505 Gln Asp Tyr Tyr Tyr Gln Ser Ala Asn Arg Ala Tyr Ser Ser Lys Thr Pro Pro Gln Asn Asn Gly Lys Lys Thr Ser Pro Asn Gly Arg Glu Lys Asn Pro Tyr Trp Val Ser Ile Gly Arg Gly Asn Val Val Thr Arg Gln 555 Ile Cys Leu Phe Gly Asn Asn Thr Tyr Thr Asp Cys Thr Pro Arg Ser Ile Asn Gly Lys Ser Tyr Tyr Ala Ala Val Arg Asp Asn Val Arg Leu Gly Arg Trp Ala Asp Val Gly Ala Gly Leu Arg Tyr Asp Tyr Arg Ser Thr His Ser Asp Asp Gly Ser Val Ser Thr Gly Thr His Arg Thr Leu Ser Trp Asn Ala Gly Ile Val Leu Lys Pro Ala Asp Trp Leu Asp Leu 635 Thr Tyr Arg Thr Ser Thr Gly Phe Arg Leu Pro Ser Phe Ala Glu Met 645 650 Tyr Gly Trp Arg Ser Gly Asp Lys Ile Lys Ala Val Lys Ile Asp Pro 665 Glu Lys Ser Phe Asn Lys Glu Ala Gly Ile Val Phe Lys Gly Asp Phe 675 Gly Asn Leu Glu Ala Ser Trp Phe Asn Asn Ala Tyr Arg Asp Leu Ile 695 Val Arg Gly Tyr Glu Ala Gln Ile Lys Asp Gly Lys Glu Gln Val Lys 715 Gly Asn Pro Ala Tyr Leu Asn Ala Gln Ser Ala Arg Ile Thr Gly Ile

725 730 735

Asn Ile Leu Gly Lys Ile Asp Trp Asn Gly Val Trp Asp Lys Leu Pro
740 745 750

Glu Gly Trp Tyr Ser Thr Phe Ala Tyr Asn Arg Val Arg Val Arg Asp
755 760 765

Ile Lys Lys Arg Ala Asp Arg Thr Asp Ile Gln Ser His Leu Phe Asp 770 780

Ala Ile Gln Pro Ser Arg Tyr Val Val Gly Ser Gly Tyr Asp Gln Pro 785 790 795 800

Glu Gly Lys Trp Gly Val Asn Gly Met Leu Thr Tyr Ser Lys Ala Lys 805 810 815

Glu Ile Thr Glu Leu Leu Gly Ser Arg Ala Leu Leu Asn Gly Asn Ser 820 825 830

Arg Asn Thr Lys Ala Thr Ser Arg Arg Thr Arg Pro Trp Tyr Ile Val 835 840 845

Asp Val Ser Gly Tyr Tyr Thr Val Lys Lys His Phe Thr Leu Arg Ala 850 855 860

Gly Val Tyr Asn Leu Leu Asn His Arg Tyr Val Thr Trp Glu Asn Val 865 870 875 880

Arg Gln Thr Ala Ala Gly Ala Val Asn Gln His Lys Asn Val Gly Val 885 890 895

Tyr Asn Arg Tyr Ala Ala Pro Gly Arg Asn Tyr Thr Phe Ser Leu Glu
900 905 910

Met Lys Phe 915

<210> 25

<211> 657

<212> PRT

<213> Moraxella catarrhalis

<400> 25

Met Lys Ser Val Pro Leu Ile Ser Gly Gly Leu Ser Phe Leu Leu Ser 1 5 10 15

Ala Cys Ser Gly Gly Ser Phe Asp Val Asp Asn Val Ser Asn Thr

Pro Ser Ser Lys Pro Arg Tyr Gln Asp Asp Thr Ser Asn Gln Arg Lys 35 40 45

Lys Ser Asn Leu Lys Lys Leu Phe Ile Ser Leu Gly Tyr Gly Met Lys 50 55 60

Leu Val Ala Gln Asn Leu Arg Gly Asn Lys Glu Pro Ser Phe Leu Asn 65 70 75 80

Glu Asp Asp Tyr Ile Ser Tyr Phe Ser Ser Leu Ser Thr Ile Glu Lys 85 90 95

Asp Val Lys Asp Asn Lys Asn Gly Ala Asp Leu Ile Gly Ser Ile Asp 100 105 110

Glu Pro Ser Thr Thr Asn Pro Pro Glu Lys His His Gly Gln Lys Tyr 115 120 125

Val Tyr Ser Gly Leu Tyr Tyr Thr Pro Ser Trp Ser Leu Asn Asp Ser 130 135 140

Lys Asn Lys Phe Tyr Leu Gly Tyr Tyr Gly Tyr Ala Phe Tyr Tyr Gly
145 150 155 160

Asn Lys Thr Ala Thr Asn Leu Pro Val Asn Gly Val Val Lys Tyr Lys 165 170 175

Gly Thr Trp Asp Phe Ile Thr Ala Thr Lys Asn Gly Lys Arg Tyr Pro 180 185 190

Leu Leu Ser Asn Gly Gly Ser His Ala Tyr Tyr Arg Arg Ser Ala Ile 195 200 205

Pro Glu Asp Ile Asp Leu Glu Asn Asp Ser Lys Asn Gly Asp Ile Gly 210 215 220

Leu Ile Ser Glu Phe Ser Ala Asp Phe Gly Thr Lys Lys Leu Thr Gly 225 230 235 240

Gln Leu Ser Tyr Thr Lys Arg Lys Thr Asn Asn Gln Pro Tyr Glu Lys 245 250 255

Lys Lys Leu Tyr Asp Ile Asp Ala Asp Ile Tyr Ser Asn Arg Phe Arg 260 265 270

Gly Thr Val Lys Pro Thr Glu Lys Asp Ser Glu Glu His Pro Phe Thr 275 280 285

Ser Glu Gly Thr Leu Glu Gly Gly Phe Tyr Pro Asn Ala Glu Glu Leu 290 295 300

Gly Gly Lys Phe Leu Ala Thr Asp Asn Arg Val Phe Gly Val Phe Ser 305 310 315 320

Ala Lys Glu Thr Glu Glu Thr Lys Lys Glu Ala Leu Ser Lys Glu Thr 325 330 335

Leu Ile Asp Gly Lys Leu Ile Thr Phe Ser Thr Lys Lys Thr Asp Ala 340 345 350

Lys Thr Asn Ala Thr Thr Ser Thr Ala Ala Asn Thr Thr Thr Asp Thr 355 360 365

Thr Ala Asn Thr Ile Thr Asp Glu Lys Asn Phe Lys Thr Glu Asp Ile 370 . 375 380

Ser Ser Phe Gly Glu Ala Asp Tyr Leu Leu Ile Asp Lys Tyr Pro Ile 385 390 395 400

Pro Leu Leu Pro Asp Lys Asn Thr Asn Asp Phe Ile Ser Ser Lys His 405 410 415

His Thr Val Gly Asn Lys Arg Tyr Lys Val Glu Ala Cys Cys Ser Asn 420 425 430

Leu Tyr Val Lys Phe Gly Met Tyr Tyr Glu Asp Pro Leu Lys Glu Lys 435 440 445

Glu Thr Glu Thr Glu Thr Glu Lys Asp Lys Glu Lys Glu Lys 450 455 460

Glu Lys Asp Lys Asp Lys Glu Lys Gln Thr Ala Ala Thr Thr Asn Thr 465 470 475 480

Tyr Tyr Gln Phe Leu Leu Gly His Arg Thr Pro Lys Asp Asp Ile Pro
485 490 495

Lys Thr Gly Ser Ala Lys Tyr His Gly Ser Trp Phe Gly Tyr Ile Thr 500 505 510

Asp Gly Lys Thr Ser Tyr Ser Pro Ser Gly Asp Lys Lys Arg Asp Lys 515 520 525

Asn Ala Val Ala Glu Phe Asn Val Asp Phe Ala Glu Lys Lys Leu Thr 530 540

Gly Glu Leu Lys Arg His Asp Thr Gly Asn Pro Val Phe Ser Ile Glu 545 550 555 560

Ala Asn Phe Asn Asn Ser Ser Asn Ala Phe Thr Gly Thr Ala Thr Ala 565 570 575

Thr Asn Phe Val Ile Asp Gly Lys Asn Ser Gln Asn Lys Asn Thr Pro 580 585 590

Ile Asn Ile Thr Thr Lys Val Asn Gly Ala Phe Tyr Gly Pro Lys Ala
595 600 605

Ser Glu Leu Gly Gly Tyr Phe Thr Tyr Asn Gly Asn Ser Thr Ala Thr 610 615 620

Asn Ser Glu Ser Ser Ser Thr Val Ser Ser Ser Ser Asn Ser Lys Asn 625 630 635 640

Ala Arg Ala Ala Val Val Phe Gly Ala Arg Gln Gln Val Glu Thr Thr
645 650 655

Lys

<210> 26

<211> 601

<212> PRT

<213> Moraxella catarrhalis

<400> 26

Met Asn Asn Pro Leu Val Asn Gln Ala Ala Met Val Leu Pro Val Phe

1 5 10 15

Leu Leu Ser Ala Cys Leu Gly Gly Gly Gly Ser Phe Asp Leu Asp Ser 20 25 30

Val Glu Thr Val Gln Asp Met His Ser Lys Pro Lys Tyr Glu Asp Glu
35 40 45

Lys Ser Gln Pro Glu Ser Gln Gln Asp Val Ser Glu Asn Ser Gly Ala
50 55 60

Ala Tyr Gly Phe Ala Val Lys Leu Pro Arg Arg Asn Ala His Phe Asn 65 70 75 80

Pro Lys Tyr Lys Glu Lys His Lys Pro Leu Gly Ser Met Asp Trp Lys 85 90 95

Lys Leu Gln Arg Gly Glu Pro Asn Ser Phe Ser Glu Arg Asp Glu Leu 100 105 110

Glu Lys Lys Arg Gly Ser Ser Glu Leu Ile Glu Ser Lys Trp Glu Asp 115 120 125

Gly Gln Ser Arg Val Val Gly Tyr Thr Asn Phe Thr Tyr Val Arg Ser 130 135 140

Gly Tyr Val Tyr Leu Asn Lys Asn Asn Ile Asp Ile Lys Asn Asn Ile 145 150 155 160

Val Leu Phe Gly Pro Asp Gly Tyr Leu Tyr Tyr Lys Gly Lys Glu Pro 165 170 175

Ser Lys Glu Leu Pro Ser Glu Lys Ile Thr Tyr Lys Gly Thr Trp Asp 180 185 190

Tyr Val Thr Asp Ala Met Glu Lys Gln Arg Phe Glu Gly Leu Gly Ser 195 200 205

Ala Ala Gly Gly Asp Lys Ser Gly Ala Leu Ser Ala Leu Glu Glu Gly 210 215 220

Val Leu Arg Asn Gln Ala Glu Ala Ser Ser Gly His Thr Asp Phe Gly 225 230 235 240

Met Thr Ser Glu Phe Glu Val Asp Phe Ser Asp Lys Thr Ile Lys Gly 245 250 255

Thr Leu Tyr Arg Asn Asn Arg Ile Thr Gln Asn Asn Ser Glu Asn Lys 260 265 270

Gln Ile Lys Thr Thr Arg Tyr Thr Ile Gln Ala Thr Leu His Gly Asn 275 280 285

Arg Phe Lys Gly Lys Ala Leu Ala Ala Asp Lys Gly Ala Thr Asn Gly 290 295 300

Ser His Pro Phe Ile Ser Asp Ser Asp Ser Leu Glu Gly Gly Phe Tyr 305 310 315 320

Gly Pro Lys Gly Glu Glu Leu Ala Gly Lys Phe Leu Ser Asn Asp Asn 325 330 335

Lys Val Ala Ala Val Phe Gly Ala Lys Gln Lys Asp Lys Lys Asp Gly 340 345 350

Glu Asn Ala Ala Gly Pro Ala Thr Glu Thr Val Ile Asp Ala Tyr Arg 355 360 365

Ile Thr Gly Glu Glu Phe Lys Lys Glu Gln Ile Asp Ser Phe Gly Asp 370 380

Val Lys Lys Leu Leu Val Asp Gly Val Glu Leu Ser Leu Leu Pro Ser 385 390 395 400

Glu Gly Asn Lys Ala Ala Phe Gln His Glu Ile Glu Gln Asn Gly Val 405 410 415

Lys Ala Thr Val Cys Cys Ser Asn Leu Asp Tyr Met Ser Phe Gly Lys 420 425 430

Leu Ser Lys Glu Asn Lys Asp Asp Met Phe Leu Gln Gly Val Arg Thr 435 440 445

Pro Val Ser Asp Val Ala Ala Arg Thr Glu Ala Asn Ala Lys Tyr Arg 450 455 460

Gly Thr Trp Tyr Gly Tyr Ile Ala Asn Gly Thr Ser Trp Ser Gly Glu 465 470 475 480

Ala Ser Asn Gln Phe Thr Glu Gly Gly Asn Arg Ala Glu Phe Asp Val 485 490 495

Asp Phe Ser Thr Lys Lys Ile Ser Gly Thr Leu Thr Ala Lys Asp Arg 500 505 510

Thr Ser Pro Ala Phe Thr Ile Thr Ala Met Ile Lys Asp Asn Gly Phe 515 520 525

Ser Gly Val Ala Lys Thr Gly Glu Asn Gly Phe Ala Leu Asp Pro Gln 530 540

Asn Thr Gly Asn Ser His Tyr Thr His Ile Glu Ala Thr Val Ser Gly 545 550 555 560

Gly Phe Tyr Gly Lys Asn Ala Ile Glu Met Gly Gly Ser Phe Ser Phe 565 570 575

Pro Gly Asn Ala Pro Glu Gly Lys Gln Glu Lys Ala Ser Val Val Phe
580 585 590

Gly Ala Lys Arg Gln Gln Leu Val Gln 595 600 <211> 711

<212> PRT

<213> Moraxella catarrhalis

<400> 27

Met Asn Asn Pro Leu Val Asn Gln Ala Ala Met Val Leu Pro Val Phe

1 10 15

Leu Leu Ser Ala Cys Leu Gly Gly Gly Gly Ser Phe Asp Leu Asp Ser 20 25 30

Val Asp Thr Glu Ala Pro Arg Pro Ala Pro Lys Tyr Gln Asp Val Ser 35 40 45

Ser Glu Lys Pro Gln Ala Gln Gln Asp Gln Gly Gly Tyr Gly Phe Ala 50 55 60

Met Arg Leu Lys Arg Arg Asn Trp Tyr Pro Gly Ala Glu Glu Ser Glu 65 70 75 80

Val Lys Leu Asn Glu Ser Asp Trp Glu Ala Thr Gly Leu Pro Thr Lys 85 90 95

Pro Lys Glu Leu Pro Lys Arg Gln Lys Ser Val Ile Glu Lys Val Glu 100 105 110

Thr Asp Gly Asp Ser Asp Ile Tyr Ser Ser Pro Tyr Leu Thr Pro Ser 115 120 125

Asn His Gln Asn Gly Ser Ala Gly Asn Gly Val Asn Gln Pro Lys Asn 130 135 140

Gln Ala Thr Gly His Glu Asn Phe Gln Tyr Val Tyr Ser Gly Trp Phe 145 150 155 160

Tyr His Ala Ala Ser Glu Lys Asp Phe Ser Asn Lys Lys Ile Trp Lys 165 170 175

Ser Gly Asp Asp Gly Tyr Ile Phe Tyr His Gly Glu Lys Pro Ser Arg 180 185 190

Gln Leu Pro Ala Ser Gly Lys Val Ile Tyr Lys Gly Val Trp His Phe 195 200 205

Val Thr Asp Thr Lys Lys Gly Gln Asp Phe Arg Glu Ile Ile Gln Pro 210 215 220

Ser Lys Lys Gln Gly Asp Arg Tyr Ser Gly Phe Ser Gly Asp Gly Ser 225 230 235 240

Glu Glu Tyr Ser Asn Lys Asn Glu Ser Thr Leu Lys Asp Asp His Glu 245 250 255

Gly Tyr Gly Phe Thr Ser Asn Leu Glu Val Asp Phe Gly Asn Lys Lys 260 265 270

Leu Thr Gly Lys Leu Ile Arg Asn Asn Ala Ser Leu Asn Asn Asn Thr 275 280 285

Asn Asn Asp Lys His Thr Thr Gln Tyr Tyr Ser Leu Asp Ala Gln Ile 295 Thr Gly Gly Asn Pro Phe Asn Gly Thr Ala Thr Ala Thr Asp Lys Lys 310 315 Glu Asn Glu Thr Lys Leu His Pro Phe Val Ser Asp Ser Ser Ser Leu 325 Glu Gly Gly Phe Phe Gly Pro Gln Gly Glu Leu Gly Phe Arg Phe Leu Thr Asp Asp Gln Lys Val Ala Val Val Gly Ser Ala Lys Thr Lys Asp Lys Leu Glu Asn Gly Ala Ala Ala Ser Gly Ser Gly Ala Ala Ala Ser Gly Gly Ala Ala Gly Thr Ser Ser Glu Asn Ser Lys Leu Thr Thr 390 Val Leu Asp Ala Val Glu Leu Thr Leu Asn Asp Lys Lys Ile Lys Asn Leu Asp Asn Phe Ser Asn Ala Ala Gln Leu Val Val Asp Gly Ile Met 420 Ile Pro Leu Leu Pro Lys Asp Ser Glu Ser Gly Asn Thr Gln Ala Asp Lys Gly Lys Asn Gly Gly Thr Glu Phe Thr Arg Lys Phe Glu His Thr Pro Glu Ser Asp Lys Lys Asp Ala Gln Ala Gly Thr Gln Thr Asn Gly Ala Gln Thr Ala Ser Asn Thr Ala Gly Asp Thr Asn Gly Lys Thr Lys Thr Tyr Glu Val Glu Val Cys Cys Ser Asn Leu Asn Tyr Leu Lys Tyr Gly Met Leu Thr Arg Lys Asn Ser Lys Ser Ala Met Gln Ala Gly Gly 520 Asn Ser Ser Gln Ala Asp Ala Lys Thr Glu Gln Val Glu Gln Ser Met 535 Phe Leu Gln Gly Glu Arg Thr Asp Glu Lys Glu Ile Pro Thr Asp Gln 555 Asn Val Val Tyr Arg Gly Ser Trp Tyr Gly His Ile Ala Asn Gly Thr 565 Ser Trp Ser Gly Asn Ala Ser Asp Lys Glu Gly Gly Asn Arg Ala Asp Phe Thr Ile Asn Phe Ala Asp Lys Lys Ile Thr Gly Lys Leu Thr Ala

Glu Asn Arg Thr Ala Gln Thr Phe Thr Ile Glu Gly Met Ile Gln Gly 610 620

Asn Gly Phe Glu Gly Thr Ala Lys Thr Ala Glu Ser Gly Phe Asp Leu 625 630 635 640

Asp Gln Lys Asn Thr Thr Arg Thr Pro Lys Ala Tyr Ile Thr Asp Ala 645 650 655

Lys Val Lys Gly Gly Phe Tyr Gly Pro Lys Ala Glu Glu Leu Gly Gly 660 665 670

Trp Phe Ala Tyr Pro Gly Asp Lys Gln Thr Glu Lys Ala Thr Ala Thr 675 680 685

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<213> Moraxella catarrhalis

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Ser Glu Lys Pro Gln Ala Gln Lys Asp Gln Gly Gly Tyr Gly Phe Ala 50 55 60

Met Arg Phe Lys Arg Arg Asn Trp His Pro Ser Ala Asn Pro Lys Glu 65 70 75 80

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Thr Glu Pro Lys Lys Leu Pro Leu Lys Gln Gln Ser Val Ile Ser Glu
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Tyr Lys His Ala Lys Ser Glu Val Lys Asn Glu Asn Gly Leu Val Ser 170 Ala Lys Arg Gly Asp Asp Gly Tyr Ile Phe Tyr His Gly Asp Lys Pro 185 Ser Arg Gln Leu Pro Ala Ser Glu Ala Val Thr Tyr Lys Gly Val Trp 200 His Phe Val Thr Asp Thr Lys Gln Gly Gln Lys Phe Asn Asp Ile Leu Glu Thr Ser Lys Gly Gln Gly Asp Lys Tyr Ser Gly Phe Ser Gly Asp Glu Gly Glu Thr Thr Ser Asn Arg Thr Asp Ser Asn Leu Asn Asp Lys 250 His Glu Gly Tyr Gly Phe Thr Ser Asn Phe Lys Val Asp Phe Asn Asn 260 265 Lys Lys Leu Thr Gly Lys Leu Ile Arg Asn Asn Lys Val Ile Asn Thr 280 Ala Ala Ser Asp Gly Tyr Thr Thr Glu Tyr Tyr Ser Leu Asp Ala Thr 290 295 Leu Arg Gly Asn Arg Phe Ser Gly Lys Ala Ile Ala Thr Asp Lys Pro Asn Thr Gly Gly Thr Lys Leu His Pro Phe Val Phe Asp Ser Ser Ser 330 Leu Ser Gly Gly Phe Phe Gly Pro Gln Gly Glu Leu Gly Phe Arg Phe Leu Ser Asp Asp Gly Lys Val Ala Val Val Gly Ser Ala Lys Thr 360 Lys Asp Ser Thr Ala Asn Gly Asn Ala Pro Ala Ala Ser Ser Gly Pro 375 Gly Ala Ala Thr Met Pro Ser Glu Thr Arg Leu Thr Thr Val Leu Asp Ala Val Glu Leu Thr Pro Asp Gly Lys Glu Ile Lys Asn Leu Asp Asn 405 410 Phe Ser Asn Ala Thr Arg Leu Val Val Asp Gly Ile Met Ile Pro Leu 425 Leu Pro Thr Glu Ser Gly Asn Gly Gln Ala Asp Lys Gly Lys Asn Gly 435 440 Gly Thr Asp Phe Thr Tyr Glu Thr Thr Tyr Thr Pro Glu Ser Asp Lys Lys Asp Thr Lys Ala Gln Thr Gly Ala Gly Gly Met Gln Thr Ala Ser 470 475

Gly Thr Ala Thr Val Asn Gly Gly Gln Val Gly Thr Lys Thr Tyr Lys Val Gln Val Cys Cys Ser Asn Leu Asn Tyr Leu Lys Tyr Gly Leu Leu 505 Thr Arg Glu Asn Asn Asn Ser Val Met Gln Ala Val Lys Asn Ser Ser 520 Gln Ala Asp Ala Lys Thr Lys Gln Ile Glu Gln Ser Met Phe Leu Gln 535 Gly Glu Arg Thr Asp Glu Asn Lys Ile Pro Gln Glu Gln Gly Ile Val 555 Tyr Arg Gly Phe Trp Tyr Gly Arg Ile Ala Asn Gly Thr Ser Trp Ser 565 570 Gly Lys Ala Ser Asn Ala Thr Asp Gly Asn Arg Ala Lys Phe Thr Val 585 Asn Gly Asp Arg Lys Glu Ile Thr Gly Thr Leu Thr Ala Glu Asn Arg 595 600 Ser Glu Ala Thr Phe Thr Ile Asp Ala Met Ile Glu Gly Asn Gly Phe 615 Lys Gly Thr Ala Lys Thr Gly Asn Asp Gly Phe Ala Pro Asp Gln Asn 625 630 Asn Ser Thr Val Thr His Lys Val His Ile Ala Asn Ala Glu Val Gln Gly Gly Phe Tyr Gly Pro Asn Ala Glu Glu Leu Gly Gly Trp Phe Ala 665 Tyr Pro Gly Asn Glu Gln Thr Lys Asn Ala Thr Val Glu Ser Gly Asn 680 Gly Asn Ser Ala Ser Ser Ala Thr Val Val Phe Gly Ala Lys Arg Gln 695 Lys Leu Val Lys 705 <210> 29 <211> 280 <212> DNA <213> Moraxella catarrhalis

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54

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<sup>&</sup>lt;212> PRT

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Thr Ser Ser Gly Thr Gly Ser Ala Ser Thr Ser Glu Pro Lys Tyr Gln 65 70 75 80

Asp Val Pro Thr Thr Pro Asn Asn Lys Glu Gln Val Ser Ser Ile Gln
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Glu Pro Ala Met Gly Tyr Gly Met Ala Leu Ser Lys Ile Asn Leu Tyr 100 105 110

Asp Gln Gln Asp Thr Pro Leu Asp Ala Lys Asn Ile Ile Thr Leu Asp 115 120 125

Gly Lys Lys Gln Val Ala Asp Asn Gln Lys Ser Pro Leu Pro Phe Ser 130 135 140

Glu Ala Asp Lys Asn Ala Ile Gly Glu Arg Ile Lys Arg Glu Asn Glu 165 170 175

Gln Asn Lys Lys Ile Ser Asp Glu Glu Leu Ala Lys Lys Ile Lys Glu 180 185 190

Asn Val Arg Lys Ser Pro Glu Phe Gln Gln Val Leu Ser Ser Ile Lys 195 200 205

Ala Lys Thr Phe His Ser Asn Asp Lys Thr Thr Lys Ala Thr Thr Arg 210 215 220

Asp Leu Lys Tyr Val Asp Tyr Gly Tyr Tyr Leu Val Asn Asp Ala Asn 225 230 235 240

Tyr Leu Thr Val Lys Thr Asp Lys Pro Lys Leu Trp Asn Ser Gly Pro 245 250 255

Val Gly Gly Val Phe Tyr Asn Gly Ser Thr Thr Ala Lys Glu Leu Pro 260 265 270

Thr Gln Asp Ala Val Lys Tyr Lys Gly His Trp Asp Phe Met Thr Asp 275 280 285

Val Ala Lys Lys Arg Asn Arg Phe Ser Glu Val Lys Glu Thr Tyr Gln 290 295 300

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          Tyr Lys Ile Asp Ser Ser Ser Thr Gly Lys Ser Ile Val Ile Lys Asp
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          Ala Val Val Thr Gly Gly Phe Tyr Gly Pro Asn Ala Thr Glu Met Gly
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